

CRF Errors Edited by the STIC Systems Branch

Serial Number:

10/583,191

CRF Edit Date:

07/06/2006

Edited by:

DA

— Realigned nucleic acid/amino acid numbers/text in cases where the sequence text "wrapped" to the next line

— Corrected the SEQ ID NO. Sequence numbers edited were:

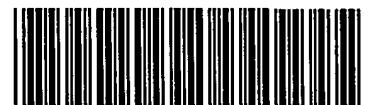
— Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

~~— Deleted: _____ invalid beginning/end-of-file text ; _____ page numbers~~

— Inserted mandatory headings/numeric identifiers, specifically:

— Moved responses to same line as heading/numeric identifier, specifically:

— Other:



IFWP

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/583,191

DATE: 07/06/2006
TIME: 13:00:24

Input Set : N:\DA\pto.da.txt
Output Set: N:\CRF4\07052006\J583191.raw

3 <110> APPLICANT: BASF Aktiengesellschaft
5 <120> TITLE OF INVENTION: Psod-Expressionseinheiten
7 <130> FILE REFERENCE: PF 55185/Mec
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/583,191
C--> 10 <141> CURRENT FILING DATE: 2006-06-14
12 <160> NUMBER OF SEQ ID NOS: 44
14 <210> SEQ ID NO: 1
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43 aagcgccatc tgacggattt tcaaaagatgt tatatgctcg gtgcggaaac ctacgaaagg 180
45 attttttacc c 191
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61 aaagatagca ctgatcggtc tgcgttagat gctgcgttag agagagcaaa aagagctgcc 180
63 gcagttgata ccaatccat agaaggaaatc ttccaaactg atcgcgggtt taccataca 240
65 gttgcaacgc aggttagggc ttgggagcaa caaatggcga tgaaaggcaa acatgttaag 300
67 cctgcgtttg acgataactct agaaggcttt gagttatgttc tcgatgcagt aactggtaga 360
69 actccaatct ctcagcaatg gattagaaat ttgcacgccc tcattctgcg gagccaagaa 420
71 agccacgagg tttttacagc cgttggagtc caaaatcagg cgcttcagaa agggcagttt 480
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/583,191

DATE: 07/06/2006
TIME: 13:00:24

Input Set : N:\DA\pto.da.txt
Output Set: N:\CRF4\07052006\J583191.raw

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 83 gcagcggaca agaataaccc gctcctgctg attagattct ttgctgaacg agtgaccgat 840
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 87 aagcttccgg atgcgctacg ccccactcgc gtattaccag aattacatga tgctcacat 960
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 93 ggctataacg cttccctga tagctattgt ctgaccctag ctttcaatag caactctcca 1140
 95 aaacaaaatct tccaccgct atccatagta atagcagctc gagatggaa aagagcgagc 1200
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 99 gaggctgttg ttactgaaag ctttcgagaa cgtgtaaaaa ttacgcccga cgggattgt 1320
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 115 20 25 30
 117 Ser Ser Arg His Phe Ser Asp Leu Lys Asp Ser Thr Asp Arg Ser Ala
 118 35 40 45
 120 Leu Asp Ala Ala Leu Glu Arg Ala Lys Arg Ala Ala Ala Val Asp Thr
 121 50 55 60
 123 Asn Ala Ile Glu Gly Ile Phe Gln Thr Asp Arg Gly Phe Thr His Thr
 124 65 70 75 80
 126 Val Ala Thr Gln Val Gly Ala Trp Glu Gln Gln Met Ala Met Lys Gly
 127 85 90 95
 129 Lys His Val Lys Pro Ala Phe Asp Asp Thr Leu Glu Gly Phe Glu Tyr
 130 100 105 110
 132 Val Leu Asp Ala Val Thr Gly Arg Thr Pro Ile Ser Gln Gln Trp Ile
 133 115 120 125
 135 Arg Asn Leu His Ala Val Ile Leu Arg Ser Gln Glu Ser His Glu Val
 136 130 135 140
 138 Phe Thr Ala Val Gly Val Gln Asn Gln Ala Leu Gln Lys Gly Glu Tyr
 139 145 150 155 160
 141 Lys Thr Gln Pro Asn Ser Pro Gln Arg Ser Asp Gly Ser Val His Ala
 142 165 170 175
 144 Tyr Ala Pro Val Glu Asp Thr Pro Ala Glu Met Ala Arg Phe Ile Ser
 145 180 185 190
 147 Glu Leu Glu Ser Lys Glu Phe Leu Ala Ala Glu Lys Val Ile Gln Ala
 148 195 200 205
 150 Ala Tyr Ala His Tyr Ala Phe Val Cys Ile His Pro Phe Ala Asp Gly
 151 210 215 220
 153 Asn Gly Arg Val Ala Arg Ala Leu Ala Ser Val Phe Leu Tyr Lys Asp
 154 225 230 235 240

RAW SEQUENCE LISTING
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Input Set : N:\DA\pto.da.txt
Output Set: N:\CRF4\07052006\J583191.raw

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160 260 265 270
162 Phe Phe Ala Glu Arg Val Thr Asp Thr Ile Asn Ser Ile Ile Val Asp
163 275 280 285
165 Leu Thr Thr Pro Ile Ala Gly Lys Ser Gly Ser Ala Lys Leu Ser Asp
166 290 295 300
168 Ala Leu Arg Pro Thr Arg Val Leu Pro Glu Leu His Asp Ala Ala His
169 305 310 315 320
171 Arg Leu Gln Glu Ser Leu Phe Thr Glu Ile Arg Ser Arg Leu Asp Glu
172 325 330 335
174 Glu Gly Lys Arg Asn Gly Leu Glu Phe Leu Leu Gln Arg Ile Phe Ile
175 340 345 350
177 Gly Ser Pro Phe Asn Leu Pro Glu Gly Tyr Asn Ala Phe Pro Asp Ser
178 355 360 365
180 Tyr Cys Leu Thr Leu Ala Phe Asn Ser Asn Ser Pro Lys Gln Ile Phe
181 370 375 380
183 His Pro Leu Ser Ile Val Ile Ala Ala Arg Asp Gly Lys Arg Ala Ser
184 385 390 395 400
186 Ser Asp Leu Val Ala Ala Thr Ser Ile Gly Tyr Asn Phe His Ala Tyr
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217 <223> OTHER INFORMATION: SEQ_ID_6
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228 <220> FEATURE:

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/583,191

DATE: 07/06/2006
TIME: 13:00:24

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Output Set: N:\CRF4\07052006\J583191.raw

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/583,191

DATE: 07/06/2006
TIME: 13:00:24

Input Set : N:\DA\pto.da.txt
Output Set: N:\CRF4\07052006\J583191.raw

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 397 gcataaaaatgcgtt gtttgcgtt tcacaatttgcgtt gaaatatca taatatctca tttactaaa 4260
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 405 <211> LENGTH: 5860
 406 <212> TYPE: DNA

VERIFICATION SUMMARY
PATENT APPLICATION: US/10/583,191

DATE: 07/06/2006
TIME: 13:00:25

Input Set : N:\DA\pto.da.txt
Output Set: N:\CRF4\07052006\J583191.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application Number
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date



IFWP

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/583,191

DATE: 06/27/2006
TIME: 10:38:45

Input Set : F:\seqlist.txt
Output Set: N:\CRF4\06272006\J583191.raw

3 <110> APPLICANT: BASF Aktiengesellschaft
5 <120> TITLE OF INVENTION: Psod-Expressionseinheiten
7 <130> FILE REFERENCE: PF 55185/Mec
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/583,191
C--> 10 <141> CURRENT FILING DATE: 2006-06-14
12 <160> NUMBER OF SEQ ID NOS: 44

ERRORED SEQUENCES

2623 <210> SEQ ID NO: 44
2624 <211> LENGTH: 7
2625 <212> TYPE: DNA
2626 <213> ORGANISM: Corynebacterium glutamicum
2628 <220> FEATURE:
2629 <223> OTHER INFORMATION: RIBOSOMALE_BINDUNGSSTELLE\SHINE-DALGARNO
2631 <400> SEQUENCE: 44
2632 qaaaqqa

E--> 2636 PF 55185
E--> 2638 42
E--> 2642 PF 55185

Does Not Comply
Corrected Diskette Needed
(pg-1)

7

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VERIFICATION SUMMARY
PATENT APPLICATION: US/10/583,191

DATE: 06/27/2006
TIME: 10:38:47

Input Set : F:\seqlist.txt
Output Set: N:\CRF4\06272006\J583191.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application Number
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:2636 M:254 E: No. of Bases conflict, LENGTH:Input:55185 Counted:9 SEQ:44
L:2636 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
M:254 Repeated in SeqNo=44
L:2642 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:2642 M:252 E: No. of Seq. differs, <211> LENGTH:Input:7 Found:11 SEQ:44